

GenCore version 5.1.3
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OM protein - protein search, using sw model

Run on:

November 27, 2002, 05:38:32 ; Search time 58.521 Seconds

(without alignments)
 241.342 Million cell updates/sec

Title: US-09-893-615-89

Sequence: 1 QIVLUSQSPAILSASPGERKTV.....CQWSSSNPPFGGTMLEIR 106

Scoring table: BLOSUM2 Gapop 10.0 , Gapext 0.5

Searched: 98470 seqs, 13250520 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

- 1: A_Geneseq_10102:*
- 2: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1980.DAT:*
- 3: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1982.DAT:*
- 4: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1983.DAT:*
- 5: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1984.DAT:*
- 6: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1985.DAT:*
- 7: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1986.DAT:*
- 8: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1987.DAT:*
- 9: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1988.DAT:*
- 10: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1989.DAT:*
- 11: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1990.DAT:*
- 12: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1991.DAT:*
- 13: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1992.DAT:*
- 14: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1993.DAT:*
- 15: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1994.DAT:*
- 16: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1995.DAT:*
- 17: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1996.DAT:*
- 18: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1997.DAT:*
- 19: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1998.DAT:*
- 20: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA1999.DAT:*
- 21: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA2000.DAT:*
- 22: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA2001.DAT:*
- 23: /SIDS2/gcgdata/geneseq/geneseq/geneseq-emb1/AA2002.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result #
 No. Score Query Length DB ID Description

RESULT 1			
ID	AAW24532	standard; protein;	106 AA.
XX	AAW24532;		
XX			
DT	27-DEC-1997 (first entry)		
XX			
DE	Anti-Factor IX Mab chimeric light chain.		
XX			
KW	Thrombosis; therapy; Factor IX; anticoagulant; monoclonal antibody; chimeric antibody; antibody engineering; light chain.		
XX			
OS	Chimeric Mabs musculus.		
OS	Chimeric Homo sapiens.		
XX			
PN	WO9726010-A1.		
XX			
PD	24-JUL-1997.		
XX			
PF	17-JAN-1997; 97WO-US00759.		
XX			
PR	24-OCT-1996; 96US-0029119.		
PR	17-JAN-1995; 96US-0010108.		
XX			
PA	(SMIK) SMITHKLIN BECHAM CORP.		
PA	(UYVE-) UNTV VERNON & STATE AGRIC COLLEGE.		
XX			
PI	Blackburn MN, Church WR, Feuerstein GZ, Gross MS;		
PI	Nichols AJ, Padlan EA, Patel AH, Sylvester LR;		
XX			
DR	WPI; 1997-385117/35.		

Murine BC2 light c
 Murine Variable re
 A fusion of anti-c
 A fusion of anti-c
 Sequence encoded b

DR N-PSDB; AAT79900.

XX XX Inhibiting thrombosis with self-limiting antibody to coagulation
CC CC factor - avoids uncontrolled bleeding by providing only partial
PT PT inhibition

XX PS Example 7; Page 128; 15pp; English.

CC This polypeptide comprises a mouse-human chimeric antibody,
CC light chain in which the variable region is derived from mouse
CC anti-human factor IX monoclonal antibody BC2 (see AAW24531) and
CC human sequences from the immunoglobulin RF-TS3 CCL framework.
CC It can be expressed in transfected mammalian cells utilising a
CC post-thromboembolic construct (see AAT79900) obtained by PCR amplification (see
CC AAT79900) of BC2 cDNA and insertion of the PCR product into
CC F9H2R/C 1-3 cDNA (see AAT77374). Claimed anti-Factor IX chimeric
CC antibodies are useful in the treatment of thrombosis.

XX SQ Sequence 106 AA;

Query Match 94.6%; Score 526; DB 18; Length 106;
Best Local Similarity 95.3%; Pred. No. 1.8e-30; Matches 101; Conservative 1; Mismatches 4; Indels 0; Gaps 0;OY 1 QTVLSQSPAILSASPGRKVMTCRASSVNMHWYQOKPSSKPLWVATSNLASGVPAR 60
1 QTVLSQSPAILSASPGRKVMTCRASSVNMHWYQOKPSSKPLWVATSNLASGVPAR 60Db 61 FGSGSGTSYSLTISRVEDAATYYCQOQNSNPPIFGGCGMIEIK 106
61 FGSGSGTSYSLTISRVEDAATYYCQOQNSNPPIFGGCGMIEIK 106Db 61 FGSGSGTSYSLTISRVEDAATYYCQOQNSNPPIFGGCGMIEIK 106
1 QTVLSQSPAILSASPGRKVMTCRASSVNMHWYQOKPSSKPLWVATSNLASGVPAR 60OY 61 FGSGSGTSYSLTISRVEDAATYYCQOQNSNPPIFGGCGMIEIK 106
1 QTVLSQSPAILSASPGRKVMTCRASSVNMHWYQOKPSSKPLWVATSNLASGVPAR 60Db 61 FGSGSGTSYSLTISRVEDAATYYCQOQNSNPPIFGGCGMIEIK 106
1 QTVLSQSPAILSASPGRKVMTCRASSVNMHWYQOKPSSKPLWVATSNLASGVPAR 60

SQ Sequence 106 AA;

Query Match 94.6%; Score 526; DB 23; Length 106;
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1 QTVLSQSPAILSASPGRKVMTCRASSVNMHWYQOKPSSKPLWVATSNLASGVPAR 60Db 61 FGSGSGTSYSLTISRVEDAATYYCQOQNSNPPIFGGCGMIEIK 106
1 QTVLSQSPAILSASPGRKVMTCRASSVNMHWYQOKPSSKPLWVATSNLASGVPAR 60OY 61 FGSGSGTSYSLTISRVEDAATYYCQOQNSNPPIFGGCGMIEIK 106
1 QTVLSQSPAILSASPGRKVMTCRASSVNMHWYQOKPSSKPLWVATSNLASGVPAR 60Db 61 FGSGSGTSYSLTISRVEDAATYYCQOQNSNPPIFGGCGMIEIK 106
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1 QTVLSQSPAILSASPGRKVMTCRASSVNMHWYQOKPSSKPLWVATSNLASGVPAR 60

SQ Sequence 106 AA;

CC The invention relates to a method for treating an animal

CC post-thromboembolic induced ischaemia or reducing required dose of a

CC thrombolytic agent in treatment of an animal post-thromboembolic induced

CC ischaemia, comprising administering an anti-factor IX antibody or its

CC fragment, optionally in combination with a plasminogen activator or

CC thrombolytic agent. The method is useful for treating

CC post-thromboembolic-induced ischaemia, for preventing a required dose of a thrombolytic

CC agent. Sequences AAU0972-AAU81004 represent antibodies and vector

CC polyptides used in the method of the invention.